

SEQUENCE LISTING

<110> Miller, Duane D.

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<120> LPA RECEPTOR AGONISTS AND ANTAGONISTS AND METHODS OF
USE

<130> 20609/181

<140>

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<150> 60/190,370

<151> 2000-03-17

<160> 26

<170> PatentIn Ver. 2.1

<210> 1

<211> 1095

<212> DNA

<213> Homo sapiens

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Phe Tyr Asn Arg Ser Gly Lys His Leu Ala Thr Glu Trp Asn Thr Val
35 40 45

Ser Lys Leu Val Met Gly Leu Gly Ile Thr Val Cys Ile Phe Ile Met
50 55 60

Leu Ala Asn Leu Leu Val Met Val Ala Ile Tyr Val Asn Arg Arg Phe
65 70 75 80

His Phe Pro Ile Tyr Tyr Leu Met Ala Asn Leu Ala Ala Asp Phe
85 90 95

Phe Ala Gly Leu Ala Tyr Phe Tyr Leu Met Phe Asn Thr Gly Pro Asn
100 105 110

Thr Arg Arg Leu Thr Val Ser Thr Trp Leu Leu Arg Gln Gly Leu Ile
115 120 125

Asp Thr Ser Leu Thr Ala Ser Val Ala Asn Leu Ala Ile Ala Ile
130 135 140

Glu Arg His Ile Thr Val Phe Arg Met Gln Leu His Thr Arg Met Ser
145 150 155 160

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165 170 175

Val Met Gly Ala Ile Pro Ser Val Gly Trp Asn Cys Ile Cys Asp Ile
180 185 190

Glu Asn Cys Ser Asn Met Ala Pro Leu Tyr Ser Asp Ser Tyr Leu Val
195 200 205

Phe Trp Ala Ile Phe Asn Leu Val Thr Phe Val Val Met Val Val Leu
210 215 220

Tyr Ala His Ile Phe Gly Tyr Val Arg Gln Arg Thr Met Arg Met Ser
225 230 235 240

Arg His Ser Ser Gly Pro Arg Arg Asn Arg Asp Thr Met Met Ser Leu
245 250 255

Leu Lys Thr Val Val Ile Val Leu Gly Ala Phe Ile Ile Cys Trp Thr
260 265 270

Pro Gly Leu Val Leu Leu Leu Asp Val Cys Cys Pro Gln Cys Asp
275 280 285

Val Leu Ala Tyr Glu Lys Phe Phe Leu Leu Leu Ala Glu Phe Asn Ser
290 295 300

Ala Met Asn Pro Ile Ile Tyr Ser Tyr Arg Asp Lys Glu Met Ser Ala
305 310 315 320

Thr Phe Arg Gln Ile Leu Cys Cys Gln Arg Ser Glu Asn Pro Thr Gly
325 330 335

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<213> Homo sapiens

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25

30

Val	Val	Val	Val	Ala	Leu	Gly	Leu	Thr	Val	Ser	Val	Leu	Val	Leu	Leu

35

40

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Thr	Asn	Leu	Leu	Val	Ile	Ala	Ala	Ile	Ala	Ser	Asn	Arg	Arg	Phe	His

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Gln	Pro	Ile	Tyr	Tyr	Leu	Leu	Gly	Asn	Leu	Ala	Ala	Ala	Asp	Leu	Phe

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Ala	Gly	Val	Ala	Tyr	Leu	Phe	Leu	Met	Phe	His	Thr	Gly	Pro	Arg	Thr

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Ala	Arg	Leu	Ser	Leu	Glu	Gly	Trp	Phe	Leu	Arg	Gln	Gly	Leu	Leu	Asp

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105

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Thr	Ser	Leu	Thr	Ala	Ser	Val	Ala	Thr	Leu	Leu	Ala	Ile	Ala	Val	Glu

115

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Arg	His	Arg	Ser	Val	Met	Ala	Val	Gln	Leu	His	Ser	Arg	Leu	Pro	Arg

130

135

140

Gly Arg Val Val Met Leu Ile Val Gly Val Trp Val Ala Ala Leu Gly
145 150 155 160

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165 170 175

Arg Cys Ser Arg Met Ala Pro Leu Leu Ser Arg Ser Tyr Leu Ala Val
180 185 190

Trp Ala Leu Ser Ser Leu Leu Val Phe Leu Leu Met Val Ala Val Tyr
195 200 205

Thr Arg Ile Phe Phe Tyr Val Arg Arg Arg Val Gln Arg Met Ala Glu
210 215 220

His Val Ser Cys His Pro Arg Tyr Arg Glu Thr Thr Leu Ser Leu Val
225 230 235 240

Lys Thr Val Val Ile Ile Leu Gly Ala Phe Val Val Cys Trp Thr Pro
245 250 255

Gly Gln Val Val Leu Leu Leu Asp Gly Leu Gly Cys Glu Ser Cys Asn
260 265 270

Val Leu Ala Val Glu Lys Tyr Phe Leu Leu Leu Ala Glu Ala Asn Ser
275 280 285

Leu Val Asn Ala Ala Val Tyr Ser Cys Arg Asp Ala Glu Met Arg Arg
290 295 300

Thr Phe Arg Arg Leu Leu Cys Cys Ala Cys Leu Arg Gln Ser Thr Arg
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<212> DNA

<213> Homo sapiens

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130 135 140

Val Thr Leu Leu Ile Leu Leu Val Trp Ala Ile Ala Ile Phe Met Gly
145 150 155 160

Ala Val Pro Thr Leu Gly Trp Asn Cys Leu Cys Asn Ile Ser Ala Cys
165 170 175

Ser Ser Leu Ala Pro Ile Tyr Ser Arg Ser Tyr Leu Val Phe Trp Thr
180 185 190

Val Ser Asn Leu Met Ala Phe Leu Ile Met Val Val Val Tyr Leu Arg
195 200 205

Ile Tyr Val Tyr Val Lys Arg Lys Thr Asn Val Leu Ser Pro His Thr
210 215 220

Ser Gly Ser Ile Ser Arg Arg Arg Thr Pro Met Lys Leu Met Lys Thr
225 230 235 240

Val Met Thr Val Leu Gly Ala Phe Val Val Cys Trp Thr Pro Gly Leu
245 250 255

Val Val Leu Leu Leu Asp Gly Leu Asn Cys Arg Gln Cys Gly Val Gln
260 265 270

His Val Lys Arg Trp Phe Leu Leu Leu Ala Leu Leu Asn Ser Val Val
275 280 285

Asn Pro Ile Ile Tyr Ser Tyr Lys Asp Glu Asp Met Tyr Gly Thr Met
290 295 300

Lys Lys Met Ile Cys Cys Phe Ser Gln Glu Asn Pro Glu Arg Arg Pro
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<212> DNA

<213> Homo sapiens

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<213> Homo sapiens

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25

30

Pro Pro Phe Gln His Pro Asp Leu Ser Pro Leu Leu Arg Tyr Ser Phe

35

40

45

Glu Thr Met Ala Pro Thr Gly Leu Ser Ser Leu Thr Val Asn Ser Thr

50

55

60

Ala Val Pro Thr Thr Pro Ala Ala Phe Lys Ser Leu Asn Leu Pro Leu

65

70

75

80

Gln	Ile	Thr	Leu	Ser	Ala	Ile	Met	Ile	Phe	Ile	Leu	Phe	Val	Ser	Phe
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Leu	Gly	Asn	Leu	Val	Val	Cys	Leu	Met	Val	Tyr	Gln	Lys	Ala	Ala	Met
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Arg	Ser	Ala	Ile	Asn	Ile	Leu	Leu	Ala	Ser	Leu	Ala	Phe	Ala	Asp	Met
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Phe	Trp	Leu	Phe	Val	Ile	Glu	Gly	Val	Ala	Ile	Leu	Leu	Ile	Ile	Ser
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Ala	Tyr	Val	Ile	Leu	Ile	Ser	Leu	Ile	Ser	Phe	Phe	Ile	Pro	Phe	Leu
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Ala	Leu	Arg	Ile	His	Ser	Tyr	Pro	Glu	Gly	Ile	Cys	Leu	Ser	Gln	Ala
						275			280			285			
Ser	Lys	Leu	Gly	Leu	Met	Ser	Leu	Gln	Arg	Pro	Phe	Gln	Met	Ser	Ile
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Asp	Met	Gly	Phe	Lys	Thr	Arg	Ala	Phe	Thr	Thr	Ile	Leu	Ile	Leu	Phe
						305		310			315			320	
Ala	Val	Phe	Ile	Val	Cys	Trp	Ala	Pro	Phe	Thr	Thr	Tyr	Ser	Leu	Val
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Ala Thr Phe Ser Lys His Phe Tyr Tyr Gln His Asn Phe Phe Glu Ile
340 345 350

Ser Thr Trp Leu Leu Trp Leu Cys Tyr Leu Lys Ser Ala Leu Asn Pro
355 360 365

Leu Ile Tyr Tyr Trp Arg Ile Lys Lys Phe His Asp Ala Cys Leu Asp
370 375 380

Met Met Pro Lys Ser Phe Lys Phe Leu Pro Gln Leu Pro Gly His Thr
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<212> DNA

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<210> 10

<211> 20

<212> DNA

<213> Artificial Sequence

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<210> 11

<211> 21

<212> DNA

<213> Artificial Sequence

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<210> 12
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<212> DNA
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<223> Description of Artificial Sequence: primer,
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<210> 13
<211> 22
<212> DNA
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<210> 14
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<210> 15
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<210> 16
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<210> 17
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<210> 18
<211> 21
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<210> 19
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<210> 20
<211> 22
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<400> 20
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<210> 21
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<210> 22
<211> 22
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<210> 23
<211> 22
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<210> 24
<211> 23
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reverse EDG-8

<400> 24
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<210> 25
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<220>
<223> Description of Artificial Sequence: primer,
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<400> 25
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<210> 26
<211> 22
<212> DNA
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<220>
<223> Description of Artificial Sequence: primer,
reverse PSP24

<400> 26
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22



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